

INPUT SET: S2909.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/027,205DATE: 03/07/98
TIME: 16:19:47

47 (i) SEQUENCE CHARACTERISTICS:
 48 (A) LENGTH: 7 amino acids
 49 (B) TYPE: amino acid
 50 (D) TOPOLOGY: linear

52 (ii) MOLECULE TYPE: peptide

54 (v) FRAGMENT TYPE: internal

57 (ix) FEATURE:
 58 (A) NAME/KEY: Peptide
 59 (B) LOCATION: 1
 60 (D) OTHER INFORMATION: /note= "Xaa is bewteen 0 and 20
 61 amino acids"

63 (ix) FEATURE:
 64 (A) NAME/KEY: Peptide
 65 (B) LOCATION: 7
 66 (D) OTHER INFORMATION: /note= "Xaa is between 0 and 20
 67 amino acids"

70 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

72 Xaa Pro Pro Tyr Tyr Leu Xaa
 73 1 5

75 (2) INFORMATION FOR SEQ ID NO:2:

77 (i) SEQUENCE CHARACTERISTICS:
 78 (A) LENGTH: 39 base pairs
 79 (B) TYPE: nucleic acid
 80 (C) STRANDEDNESS: single
 81 (D) TOPOLOGY: linear

83 (ii) MOLECULE TYPE: cDNA

87 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

89 CATGAAGCTT CTCGAGCCGC CACCATGGCT TGCCTTGGGA

39

91 (2) INFORMATION FOR SEQ ID NO:3:

93 (i) SEQUENCE CHARACTERISTICS:
 94 (A) LENGTH: 40 base pairs
 95 (B) TYPE: nucleic acid
 96 (C) STRANDEDNESS: single
 97 (D) TOPOLOGY: linear

99 (ii) MOLECULE TYPE: cDNA

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"Xaa" can only represent a single amino acid.
 Variable length not permitted.

Show the maximum number of Xaa's
 with variable length, and state that
 some Xaa's may be missing (place
 statement in (ix) FEATURE: section).

Xaa is bewteen 0 and 20

Xaa is between 0 and 20

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100
101
102
103 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
104
105 GAGAATTCTA GACTAGCTTA AGTCAGAACATC TGGGCACGGT 40
106
107 (2) INFORMATION FOR SEQ ID NO:4:
108
109 (i) SEQUENCE CHARACTERISTICS:
110 (A) LENGTH: 21 base pairs
111 (B) TYPE: nucleic acid
112 (C) STRANDEDNESS: single
113 (D) TOPOLOGY: linear
114
115 (ii) MOLECULE TYPE: cDNA
116
117
118
119 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
120
121 GGCTAACTAG GGAACCCACT G 21
122
123 (2) INFORMATION FOR SEQ ID NO:5:
124
125 (i) SEQUENCE CHARACTERISTICS:
126 (A) LENGTH: 24 base pairs
127 (B) TYPE: nucleic acid
128 (C) STRANDEDNESS: single
129 (D) TOPOLOGY: linear
130
131 (ii) MOLECULE TYPE: cDNA
132
133
134
135 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
136
137 CTGCTAGAGA TTTTCCACAC TGAC 24
138
139 (2) INFORMATION FOR SEQ ID NO:6:
140
141 (i) SEQUENCE CHARACTERISTICS:
142 (A) LENGTH: 30 base pairs
143 (B) TYPE: nucleic acid
144 (C) STRANDEDNESS: single
145 (D) TOPOLOGY: linear
146
147 (ii) MOLECULE TYPE: cDNA
148
149
150
151 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
152

RAW SEQUENCE LISTING
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153 CCGTCTGTTG TGTGACTCTG GTAACTAGAG

154
155 (2) INFORMATION FOR SEQ ID NO:7:156 (i) SEQUENCE CHARACTERISTICS:
157 (A) LENGTH: 30 base pairs
158 (B) TYPE: nucleic acid
159 (C) STRANDEDNESS: single
160 (D) TOPOLOGY: linear
161162 (ii) MOLECULE TYPE: cDNA
163

164

165

166 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
167168 CTTGATAATC CATCTTGTTC CACCCTGTGC
169

170

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION **US/09/027,205**

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Line Error

Original Text